



# SEQUENCE LISTING

<110> NEELAM, Beena et al

<120> ISOLATED HUMAN RAS-LIKE PROTEINS,  
NUCLEIC ACID MOLECULES ENCODING THESE HUMAN RAS-LIKE  
PROTEINS, AND USES THEREOF

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<140> 09/778,963

<141> 2001-02-08

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<212> DNA

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Lys Asn Ser Tyr Arg Met Val Val Leu Gly Ala Ser Arg Val Gly Lys
      20           25           30
Ser Ser Ile Val Ser Arg Phe Leu Asn Gly Arg Phe Glu Asp Gln Tyr
      35           40           45
Thr Pro Thr Ile Glu Asp Phe His Arg Lys Val Tyr Asn Ile Arg Gly
      50           55           60
Asp Met Tyr Gln Leu Asp Ile Leu Asp Thr Ser Gly Asn His Pro Phe
65           70           75           80
Pro Ala Met Arg Arg Leu Ser Ile Leu Thr Gly Asp Val Phe Ile Leu
      85           90           95
Val Phe Ser Leu Asp Asn Arg Glu Ser Phe Asp Glu Val Lys Arg Leu
      100          105          110
Gln Lys Gln Ile Leu Glu Val Lys Ser Cys Leu Lys Asn Lys Thr Lys
      115          120          125
Glu Ala Ala Glu Leu Pro Met Val Ile Cys Gly Asn Lys Asn Asp His
      130          135          140
Gly Glu Leu Cys Arg Gln Val Pro Thr Thr Glu Ala Glu Leu Leu Val
145          150          155          160
Ser Gly Asp Glu Asn Cys Ala Tyr Phe Glu Val Ser Ala Lys Lys Asn
      165          170          175

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Thr	Asn	Val	Asp	Glu	Met	Phe	Tyr	Val	Leu	Phe	Ser	Met	Ala	Lys	Leu
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Pro	His	Glu	Met	Ser	Pro	Ala	Leu	His	Arg	Lys	Ile	Ser	Val	Gln	Tyr
		195					200					205			
Gly	Asp	Ala	Phe	His	Pro	Arg	Pro	Phe	Cys	Met	Arg	Arg	Val	Lys	Glu
	210					215					220				
Met	Asp	Ala	Tyr	Gly	Met	Val	Ser	Pro	Phe	Ala	Arg	Arg	Pro	Ser	Val
225					230					235					240
Asn	Ser	Asp	Leu	Lys	Tyr	Ile	Lys	Ala	Lys	Val	Leu	Arg	Glu	Gly	Gln
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Ala	Arg	Glu	Arg	Asp	Lys	Cys	Thr	Ile	Gln						
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Ser	Ser	Ile	Val	Ser	Arg	Phe	Leu	Asn	Gly	Arg	Phe	Glu	Asp	Gln	Tyr
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Thr	Pro	Thr	Ile	Glu	Asp	Phe	His	Arg	Lys	Val	Tyr	Asn	Ile	His	Gly
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Asp	Met	Tyr	Gln	Leu	Asp	Ile	Leu	Asp	Thr	Ser	Gly	Asn	His	Pro	Phe
65					70					75					80
Pro	Ala	Met	Arg	Arg	Leu	Ser	Ile	Leu	Thr	Gly	Asp	Val	Phe	Ile	Leu
				85						90				95	
Val	Phe	Ser	Leu	Asp	Ser	Arg	Glu	Ser	Phe	Asp	Glu	Val	Lys	Arg	Leu
			100					105					110		
Gln	Lys	Gln	Ile	Leu	Glu	Val	Lys	Ser	Cys	Leu	Lys	Asn	Lys	Thr	Lys
		115					120					125			
Glu	Ala	Ala	Glu	Leu	Pro	Met	Val	Ile	Cys	Gly	Asn	Lys	Asn	Asp	His
	130					135					140				
Ser	Glu	Leu	Cys	Arg	Gln	Val	Pro	Ala	Met	Glu	Ala	Glu	Leu	Leu	Val
145					150					155					160
Ser	Gly	Asp	Glu	Asn	Cys	Ala	Tyr	Phe	Glu	Val	Ser	Ala	Lys	Lys	Asn
				165					170					175	
Thr	Asn	Val	Asn	Glu	Met	Phe	Tyr	Val	Leu	Phe	Ser	Met	Ala	Lys	Leu
			180					185					190		
Pro	His	Glu	Met	Ser	Pro	Ala	Leu	His	His	Lys	Ile	Ser	Val	Gln	Tyr
		195					200					205			
Gly	Asp	Ala	Phe	His	Pro	Arg	Pro	Phe	Cys	Met	Arg	Arg	Thr	Lys	Val
	210					215					220				
Ala	Gly	Ala	Tyr	Gly	Met	Val	Ser	Pro	Phe	Ala	Arg	Arg	Pro	Ser	Val
225					230					235					240
Asn	Ser	Asp	Leu	Lys	Tyr	Ile	Lys	Ala	Lys	Val	Leu	Arg	Glu	Gly	Gln
			245						250					255	
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			260					265							

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